


TECH CENTER 1600/2900



TRADEMARK OFFICE

$$f_{\text{max}} = \frac{1}{2\pi} \sqrt{\frac{1}{m} \left(\frac{1}{\Delta x} \right)^2 \left(\frac{1}{\Delta t} \right)^2} = \frac{1}{2\pi} \sqrt{\frac{1}{m} \left(\frac{1}{\Delta x} \right)^2 \left(\frac{1}{\Delta t} \right)^2}$$

1. What is the purpose of the study?

Expressing Gene Expression in Plants

015129 734297, 01

40, 34, 35, 22

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1970-1971, 1972

65-15349-1-20

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41703 - WATER, IN Vol. 2. 6

220

4417 1857

Figure 1. The effect of the number of trials on the number of correct responses.

1973. *Archidopis trilineata*

• 44 •

[illegible]

2. $\int_0^1 \int_0^1 \frac{1}{1+x^2+y^2} dx dy = \frac{\pi}{2}$

• \mathbb{Z}_2 is a subring of \mathbb{Z}_4 and \mathbb{Z}_4 is a \mathbb{Z}_2 -module. \mathbb{Z}_4 is a free \mathbb{Z}_2 -module of rank 2. \mathbb{Z}_4 is not a free \mathbb{Z}_4 -module of rank 1.

[illegible][illegible]

P^2

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410a-2

Met Asp Thr Gly Gly Asn Ser Leu Ala Ser Gly Trp Asp Gly Val Lys

1

5

9

13

Arg Lys Val Cys Tyr Phe Tyr Asp Phe Ala Val Gly Asn Tyr Tyr Tyr

20

25

30

Gly Gln Gly His Pro Met Lys Pro His Arg Ile Arg Met Thr His Ala

35

40

45

Leu Leu Ala His Tyr Gly Leu Leu Gln His Met Thr Val Leu Lys Pro

50

55

60

Phe Pro Ala Arg Glu Arg Asp Leu Cys Arg Phe His Ala Asp Asp Tyr

65

70

75

80

Val Ser Phe Leu Arg Ser Ile Thr Phe Gln Thr Gln Gln Asp Gln Ile

85

90

95

Arg Gln Leu Lys Arg Phe Asn Val Gly Glu Asp Cys Pro Val Phe Asp

100

105

110

Gly Leu Tyr Ser Phe Tyr Gln Thr Tyr Ala Gly Gly Ser Val Gly Gly

115

120

125

Ser Val Lys Leu Asn His Gly Leu Tyr Arg Ile Ala Ile Asn Tyr Ala

130

135

140

Gly Tyr Leu His His Ala Lys Lys Tyr His Ala Ser Gly His Tyr Tyr
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Val Asn Asp His Val Leu Ala His Leu His Leu Leu Lys Ser His His
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Arg Val Leu Tyr Val Asp His Asp Leu His His Gly Asp Gly Val His
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Gly Ala Phe Tyr Ala Thr Asp Arg Val Met Thr Val Ser Phe His Lys
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Phe Gly Asp Tyr Phe His His Thr Gly His His His Asp His Gly Tyr
229 230 231 232 233 234 235 236 237 238 239 240

Gly Ser Gly Lys Tyr Tyr Ser Leu Asn Val Pro Leu Asp Asp Gly His
241 242 243 244 245 246 247 248 249 250 251 252

Asp Asp His Ser Tyr His Leu Leu His Lys Pro His Met Gly Lys Val
253 254 255 256 257 258 259 260 261 262 263 264

Met Gly His Phe Arg His Gly Ala Val Val Leu His Tyr Gly Ala Asp
265 266 267 268 269 270 271 272 273 274 275 276

Ser Leu Phe Gly Asn Arg Leu Gly Tyr His Asn Leu Ser His Lys Gly
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His Ala His Tyr Val Lys His Met Arg Ser His Asn Val His Leu Leu
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Leu Leu Gly Gly Gly Gly Tyr Thr Ile Arg Asn Val Ala Arg Cys Thr
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Lys Tyr His Thr Gly Val Ala Leu Gly Val Ala Val Glu Asp Lys Met
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Pro Glu His Glu Tyr Tyr Glu Tyr Phe Gly Pro Asp Tyr Thr Leu His
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Val Ala Pro Ser Asn Met Ile Asn Lys Asn Ser Arg Glu Met Leu Glu
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Glu Ile Arg Asn Asp Leu Leu His Asn Leu Ser Lys Leu Glu His Ala
392 396 400

Pro Ser Val Pro Phe Glu Glu Arg Pro Pro Asp Thr Glu Thr Pro Glu
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Val Asp Glu Asp Glu Glu Asp Gly Asp Lys Arg Trp Asp Pro Asp Ser
420 424 428

Asp Met Asp Val Asp Asp Asp Arg Lys Leu Ile Pro Ser Arg Val Lys
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Arg Glu Ala Val Glu Pro Asp Thr Lys Asp Lys Asp Gly Leu Lys Gly
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Ile Met Leu Arg Gly Lys Gly Tyr Glu Val Glu Val Asp Leu Ser Gly
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R²

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Val Lys Met Ala His His Lys Thr Asn Lys Phe Gly Ala His Lys Ala
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Phe Pro Pro Lys Thr
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4495-4

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Arg Lys Arg Arg Val Ser Tyr Pro Tyr Ala Pro Thr Ile Gly Asp Tyr

27 24 41

Tyr Tyr Gly Glu Gly His Pro Met Lys Pro His Arg Ile Arg Met Ala

35 43 45

His Ser Leu Ile Ile His Tyr His Leu His Arg Arg Leu Glu Ile Ser

56 51 66

Arg Pro Ser Leu Ala Asp Ala Ser Asp Ile Gly Arg Pro His Ser Pro

65 73 75 80

Glu Tyr Val Asp Phe Leu Ala Ser Val Ser Pro Glu Ser Met Gly Asp

81 97 95

Pro Ser Ala Ala Arg Asn Leu Arg Arg His Asn Val Gly Glu Asp Cys

109 104 110

Pro Val Phe Asp Gly Ile Phe Asp Ile Cys Arg Ala Ser Ala Gly Gly

115 123 125

Arg Ile Gly Ala Ala Val Lys Leu Asn Arg Glu Asp Ala Asp Ile Ala

14 146 14

Ile Asn Thr Lys Lys Lys Ile His His Ala Lys Tyr Ser Phe Ala Ser

14 17 17 17

Gly Phe Tyr Tyr Val Asn Asp Ile Val Leu Gly Ile Leu Glu Leu Leu

182 187 191

Lys Met Phe Lys Arg Val Lys Tyr Ile Asp Ile Asp Val His His Gly

196 198 199

Asp Gly Val Glu Glu Ala Phe Tyr Thr Thr Asp Arg Val Met Thr Val

195 200 203

Met Phe His Lys Leu Gly Asp Phe Phe Pro Gly Thr Gly His Ile Arg

210 215 220

Asp Val Gly Ala Glu Lys Gly Lys Tyr Tyr Ala Leu Asn Val Pro Leu

225 230 235 240

Asn Asp Gly Met Asp Asp Glu Ser Phe Arg Ser Leu Phe Arg Pro Leu

245 250 251

Ile Glu Lys Val Met Glu Val Tyr Glu Pro His Ala Val Val Leu Glu

255 261 270

Tyr Gly Ala Asp Ser Leu Ser Gly Asp Arg Leu Gly Tyr Phe Asn Leu

275 281 281

Asn Val Lys Gly His Ala Asp Tyr Leu Asn Ile Leu Arg Ser Tyr Asn

28 290 297

Val Leu Leu Met Val Leu Lys Gly Leu Gly Lys Thr Ile Arg Asn Val

300 301 307 307

Ala Arg Tyr Trp Tyr Tyr Glu Thr Ala Val Ala Val Gly Val Glu Trp

325

33

340

Asp Asn Lys Leu Phe Tyr Arg Glu Tyr Phe Ala Tyr Ile Gly Trp Asn

345

345

350

Tyr Thr Leu His Val Asp Pro Ser Pro Met Glu Asn Leu Asn Thr Pro

355

360

365

Lys Asp Met Glu Arg Ile Arg Asn Thr Leu Leu Glu Glu Leu Ser Gly

370

375

380

Leu Ile His Ala Pro Ser Val Glu Phe Glu His Thr Pro Pro Val Asn

385

390

395

400

Arg Val Leu Asp Glu Pro Glu Asp Asp Met Glu Thr Arg Pro Lys Pro

405

410

415

Arg Xaa Trp Ser Gly Thr Ala Thr Tyr Glu Ser Asp Ser Asp Asp Asp

420

425

430

Asp Lys Pro Leu His Gly Tyr Ser Tyr Arg Gly Gly Ala Thr Thr Asn

435

440

445

Arg Asp Ser Thr Gly Glu Asp Glu Met Asp Asn Asn Asn Pro Glu Pro

450

455

460

Asp Val Asn Leu Pro Ser Ser

465

47

Met Glu Phe Trp Gly Ile His Val Lys Ser Gly Lys Phe Val Thr Val
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Thr Pro Glu Glu Gly Ile Leu Ile His Val Ser Glu Ala Ser Leu Gly
20 25 30

Glu Cys Lys Asn Lys Lys Gly Glu Phe Val Pro Leu His Val Lys Val
35 40 45

Gly Asn Gln Asn Leu Val Leu Gly Thr Leu Ser Thr Glu Asn Ile Pro
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Gln Leu Phe Cys Asp Leu Val Phe Asp Lys Glu Phe Glu Leu Ser His
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Thr Trp Gly Lys Gly Ser Val Tyr Phe Val Gly Tyr Lys Thr Pro Asn
85 90 95

Ile Glu Pro Gln Gly Tyr Ser Glu Glu Glu Glu Glu Glu Glu Glu
100 105 110

Val Pro Ala Gly Asn Ala Ala Lys Ala Val Ala Lys Pro Lys Ala Lys
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Pro Ala Glu Val Lys Pro Ala Val Asp Asp Glu Glu Asp Glu Ser Asp
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Ser Asn Gly Ser Asp Glu Asp Asp Ser Asp Gly Glu Asp Ser Glu Glu
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20

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30

Arg Thr Val Lys Ser Gly His Ser Val Val Leu Ser Val Thr Val Gly

35

40

45

Gly Ala Lys Leu Val Ile Gly Thr Leu Ser Gln Asp Lys Phe Pro Gln

55

60

65

Ile Ser Phe Asp Leu Val Phe Asp Lys Phe Phe Gln Leu Ser His Ser

65

70

75

80

Gly Thr Lys Ala Asn Val His Phe Ile Gly Tyr Lys Ser Pro Asn Ile

85

90

95

Glu Gln Asp Asp Phe Thr Ser Ser Asp Asp Glu Asp Val Pro Glu Ala

100

105

110

Val Pro Ala Pro Ala Pro Thr Ala Val Thr Ala Asn Gly Asn Ala Gly

115

120

125

Ala Ala Val Val Lys Ala Asp Thr Lys Pro Lys Ala Lys Pro Ala Glu

130

135

140

Val Lys Pro Ala Glu Glu Lys Pro Glu Ser Asp Glu Glu Asp Glu Ser

145

150

155

160

Asp Asp Glu Asp Glu Ser Glu Glu Asp Asp Asp Ser Glu Lys Gly Met

165

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